```
==============================================================
Sequence Listing was accepted.
See attached Validation Report.
If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).
Reviewer: Keisha Douglas
Timestamp: [year=2007; month=12; day=18; hr=19; min=2; sec=7; ms=999; ]
==============================================================
```

Application No: 10562023   Version No: 1.1

                    Input Set:

                    Output Set:

            Started: 2007-12-18 19:01:20.920
           Finished: 2007-12-18 19:01:20.936
            Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 16 ms
     Total Warnings: 0
       Total Errors: 0
 No. of SeqIDs Defined: 3
   Actual SeqID Count: 3

Sequence listing

<110> Schuster, Matthias

<120> Method for investigating cytosine methylation in DNA
      sequences by means of triplex-forming oligomers

<130> 82493

<140> 10/562,023
<141> 2005-12-20

<160> 3

<210> 1
<211> 143
<212> DNA
<213> HOMO SAPIENS

<400> 1
cggaggaaga aagaggaggg gctggctggt caccagaggg tggggcggac cgcgtgcgct 60
cggcggctgc ggagaggggg agagcaggca gcgggcggcg gggagcagca tggagccggc 120
ggcggggagc agcatggagc ctt   143

<210> 2
<211> 131
<212> DNA
<213>HOMO SAPIENS

<400> 2
cggaggaaga aagaggaggg gctggctggt caccagaggg tggggcggac cgcgtgcgct 60
cggcggctgc ggagaggggg agagcaggca gcgggcggcg gggagcagca tggagccggc 120
ggcggggagc a   131

<210> 3
<211> 143
<212> DNA
<213> HOMO SAPIENS

<400> 3
cggaggaaga aagaggaggg gctggctggt caccagaggg tggggcggac cgcgtgcgct 60
cggcggctgc ggagaggggg agagcaggca gcgggcggcg gggagcagca tggagccggc 120
ggcggggagc agcatggagc ctt   143